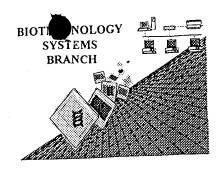
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/829, 113
Source:	OIPE
Date Processed by STIC:	4-27-01

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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: <u>patin21help@uspto.gov</u> or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: <u>patin3help@uspto.gov</u> or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

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Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/829,113

DATE: 04/27/2001 TIME: 13:07:31

Input Set : A:\209598.txt

Output Set: N:\CRF3\04272001\I829113.raw

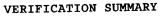
Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Evans, William McDonald, Oliver 6 < 120> TITLE OF INVENTION: HAPLOTYPING METHOD FOR MULTIPLE DISTAL NUCLEOTIDE POLYMORPHISMS)

8 < 130 > FILE REFERENCE: 44158/209598 (5853-3)

C--> 10 < 140 > CURRENT APPLICATION NUMBER: US/09/829,113 next line to correct line

C--> 10 < 141 > CURRENT RETURNS DAME: 2001-04 00 C--> 10 <141> CURRENT FILING DATE: 2001-04-09 length error. 10 <160> NUMBER OF SEQ ID NOS: 4 12 <170> SOFTWARE: PatentIn version 3.0 14 <210> SEQ ID NO: 1 Valid (213) responses are: 15 <211> LENGTH: 35 16 <212> TYPE: DNA 17 <213> ORGANISM: Oligonucleotide primer 1 - Genus species of organism 19 <220> FEATURE: 20 <221> NAME/KEY: misc_feature 21 <222> LOCATION: (3)..(8) 22 <223> OTHER INFORMATION: BamHI site 2 - Unknown 25 <400> SEQUENCE: 1 3 - Artificial sequence. 26 gcggatccga ggctgctgcc acaggctcct aaaac 29 <210> SEQ ID NO: 2 30 <211> LENGTH: 40 Unknown and artificial 31 <212> TYPE: DNA 32 <213> ORGANISM: Oligonucleotide primer) sequences must have (220),(223) 34 <220> FEATURE: 35 <221> NAME/KEY: misc_feature fectures to explain the source 36 <222> LOCATION: (3)..(8) 37 <223> OTHER INFORMATION: BamHI site of the genetic 40 material 40 <400> SEQUENCE: 2 41 geggateeca etecaggttg ggeaacaaga aegaaaetee in the sequences. 44 <210> SEQ ID NO: 3 45 <211> LENGTH: 25 46 <212> TYPE: DNA 47 <213> ORGANISM: 6ligonucleotide primer 49 <400> SEQUENCE: 3 25 50 caageettat ageettacae ceagg 53 <210> SEQ ID NO: 4 54 <211> LENGTH: 23 55 <212> TYPE: DNA 56 <213> ORGANISM: oligonucleotide primer 58 <400> SEQUENCE: 4 . 23 59 gagacagagt ttcaccatct tgg



PATENT APPLICATION: US/09/829,113

DATE: 04/27/2001 TIME: 13:07:32

Input Set : A:\209598.txt

Output Set: N:\CRF3\04272001\1829113.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date